

FIG. 1

The Genomic Structure of the Mouse *Csx/Nkx2-5*

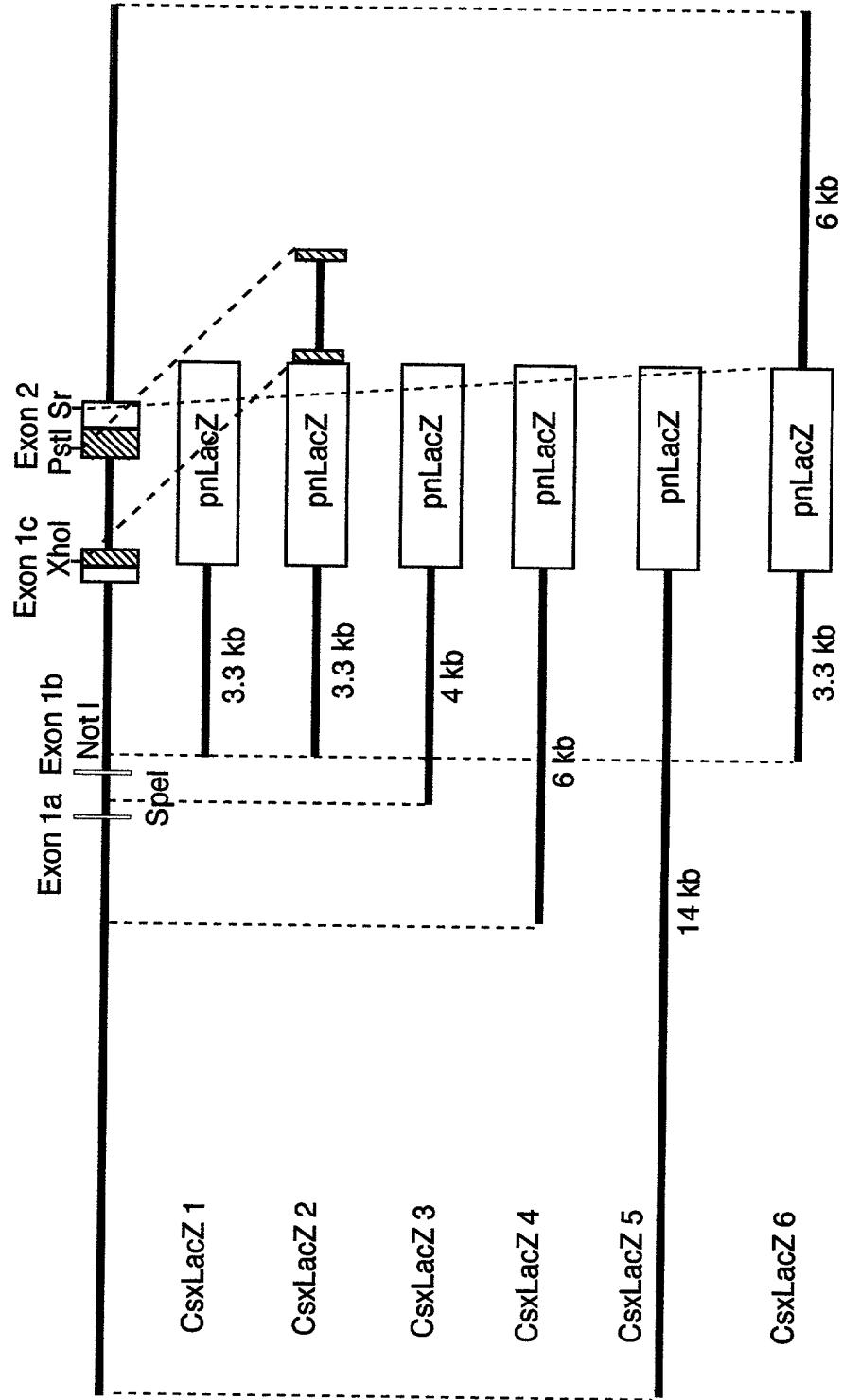


FIG. 2

The Locations of the *Csx/Nkx2.5* Cardiac Enhancers

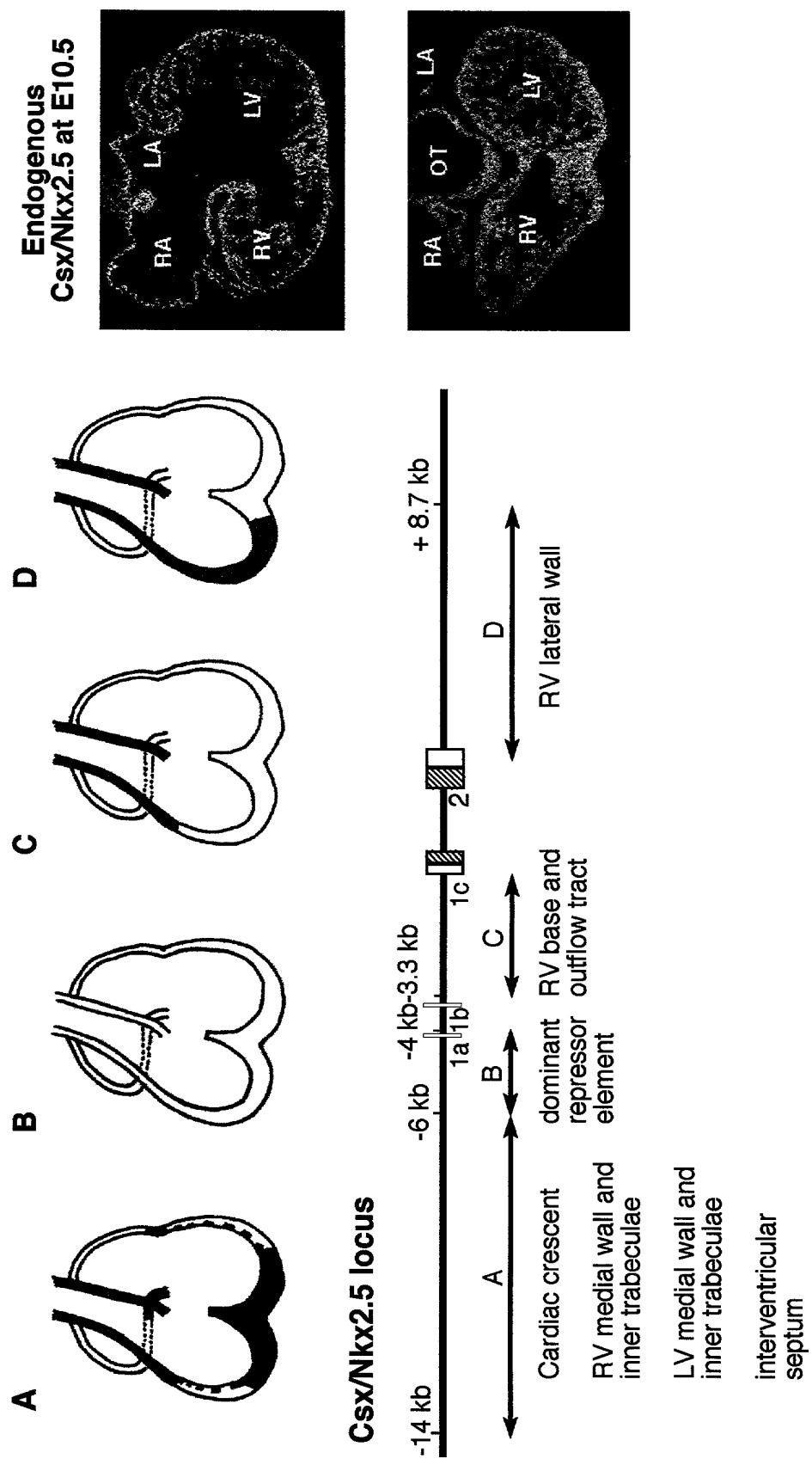


FIG. 3A

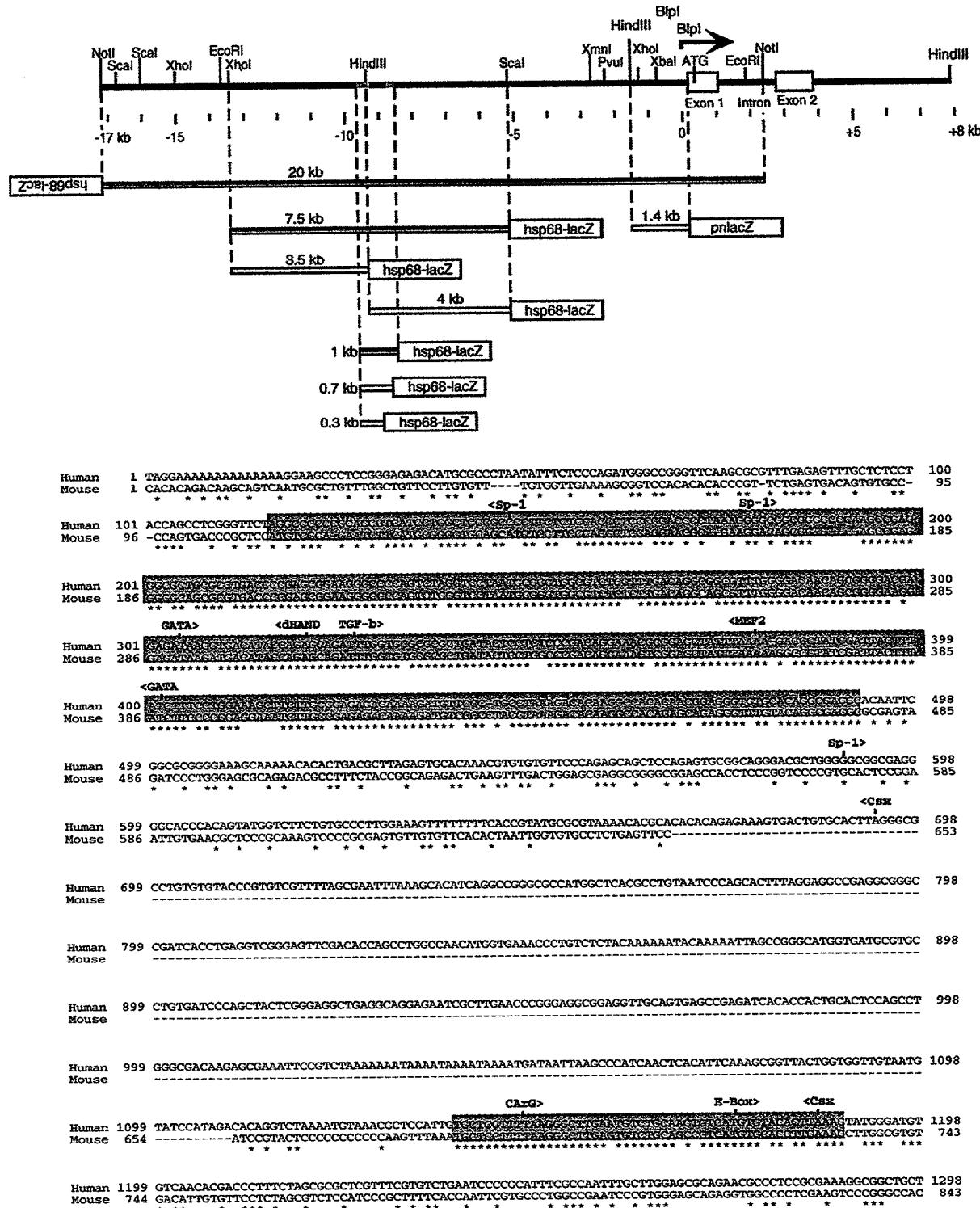
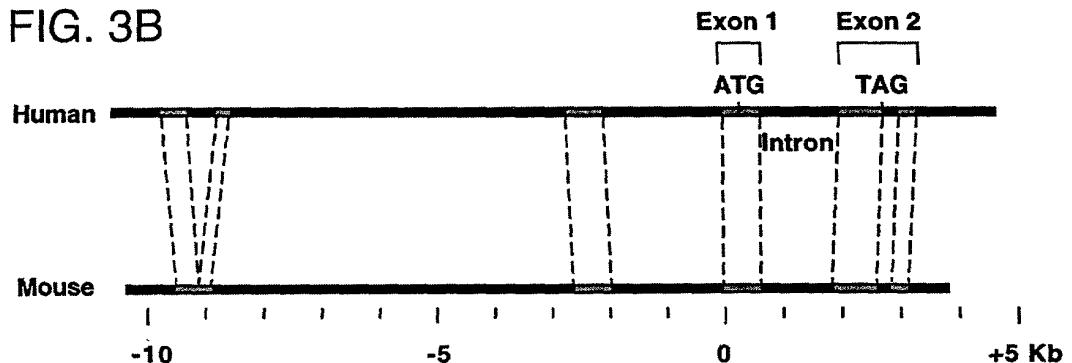


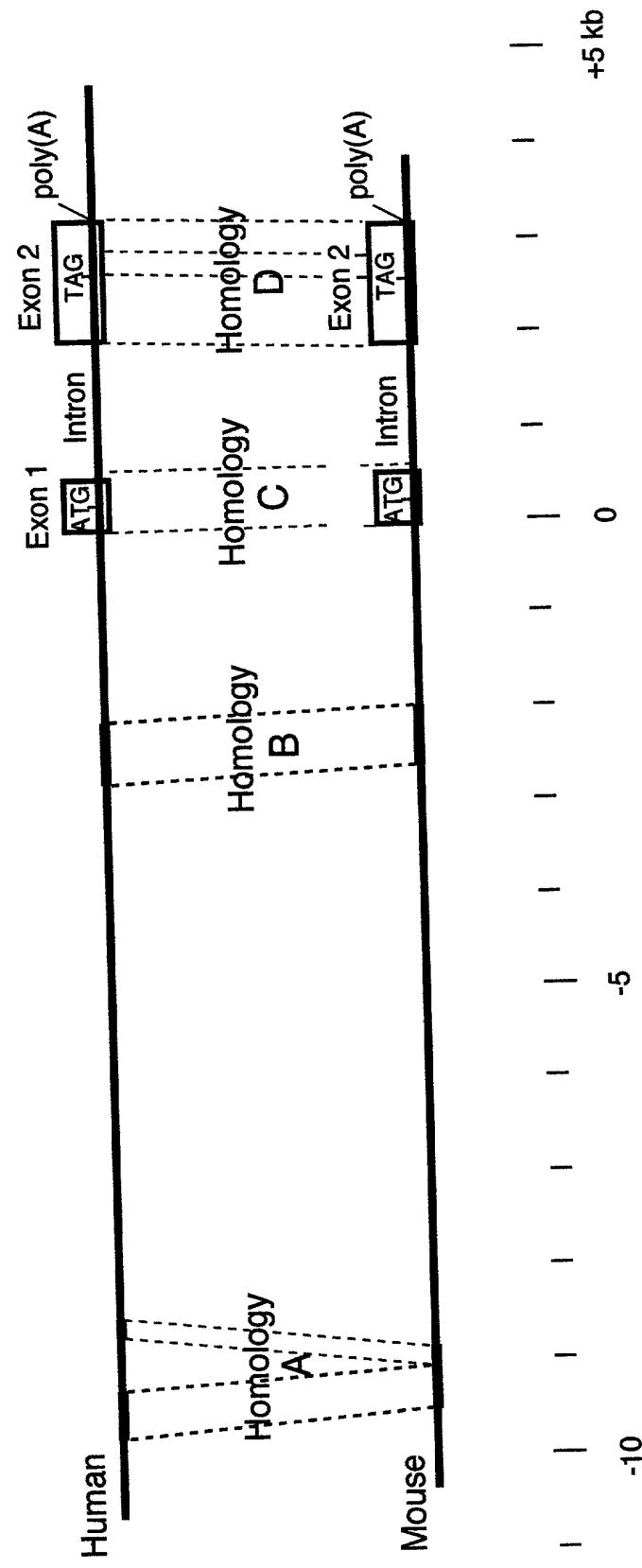
FIG. 3B



| | | | |
|-------|-----|---|------|
| Human | 1 | AGAGAACATTCATTACCGGATTACAAGAGCATAGAGACTGTAACTCACTGATCTTGTCAATTAGGGAGAGTTTTTTCCCTCCCTTTGTA | 96 |
| Mouse | 1 | GAAGAGAAAGGCCAGGGCTT-TCACCAAGGTGTAG-GGGTGGAAACATCACTGATTTGTCAAAATTAGAAGAGTTT-----CTCTCTTCTCTG | 93 |
| Human | 97 | ACACCTGACCACAGGACTGACAGTCTAGGAAGCCCCCTTACCGGAAATTAGGAAATAATTCTTGCCACCTTGATTTGCAAGGGCAATGCTAATT | 196 |
| Mouse | 94 | ACACCTGACCACAGGACTTGTCAACTCT-GGAAGCCCTTATTCGGAAA-AAGTGATAAACTCTCACCAATTCTGCAAGGGAAACG----- | 182 |
| Human | 197 | TTCTTTCTCAGAGCTCTCAAAAAAAAAAAAAACCTTACTAAAAACAGGGATCCCGGATGAGCTCTGATGTCCTCCCTAATTACCGTAATTATTC | 296 |
| Mouse | 183 | -----CCCCTCTCAAACTTATTTTTAAAGACTTAAACAGGGATCCCGGATGGGGGCCCCAATATTGCTCC-----CCCATTAAACGGTAATTATTC | 275 |
| Human | 297 | AGGCGTCCGCTCACACTAACTCTTCAAACTGTCATCGCGAGCCGCTGGGACAGATTCACTT-AACAGGCTCCGAGGCCCTGTTCCGAGCTCTT | 395 |
| Mouse | 276 | AGGCGTCACTCACACTAACTCTTCAAACTGTCATCGCGATTCGGCAGGGACAGATTCACTTAAACAGGCTCCGAGGCCCTGTTCCGAGCTCTT | 374 |
| Human | 396 | TGAGCGAGACATTAAATTGAACTGGATGTGGCTGTTGGCAGACGTACCGGCTGGGOGATAGGCATCTCTCIAACGGACACCCCCCCCCGGCGCGT | 495 |
| Mouse | 375 | TGAGCAAGACATTAAATTGAACTGGATGTGGCTGTTGGCAGACGTACCGGCTGGGOGATAGACAGATCTCGTCAACCCACACAAACAACTCTCAATCT | 474 |
| Human | 496 | CGAAAAACAACTCTCA 510 | |
| Mouse | 475 | CGGGAGGGTCCAAAGC 490 | |
| Human | 1 | TAGGAAAAAAAAAAAAAGGAAGCCCTCCGGGAGAGACATGGCCCTTAATTATTCTCCAGATGGGCCGGTTCAAGCCGTTGAGAGTTGTGCTCTCT | 100 |
| Mouse | 1 | CACACAGACAAGCACTCAATCGCTTGTGGCTGTTCTGTT-----TGTGTTGAAAGGGTCCACACACACCCCT-TCTGAGTGACAGTGCGC----- | 95 |
| Human | 101 | ACACAGGCTCGGTCTAGGCCCGACCCCTCATCCGCTCTCCCGCCCTTCCTCCACCCCTCCGGGACCCCTAAAGGGCGGGGGGGCCCAAGCCGAG | 200 |
| Mouse | 96 | -CCAGTGACCCCGCTCATGTCCTA-GAATCTTCATGGGGCTGCGCTCAGCTCTCTTCTCCACCCCTCAGGAACGC--GAAGGACAGCGG-----AGCGGAG | 185 |
| Human | 201 | GGCGCTTCGGCTGACCCCCGGAGCGGAGGGGCCCTCTCTGGCTCTCTAATCTGGGGGGCTGCGCTCTCTGACAGGGGGGTTGGGCAACACACGGGAGG | 300 |
| Mouse | 186 | GGGGCAGCGCGSTGACCCCGAGCCGGAGGGGCCCTCTCTGGCTCTCTAATCTGGGGGGCTGCGCTCTCTGACAGGGGGTTGGGCAACACACGGGAGG | 285 |
| Human | 301 | GAGAGTAAAGGTGACATACCAAGAGCAGATTTTGCGCGGGCTGTGAACTCTCTCTCCGACAGGAACCGGGAGCTTAAAGGGCTATCGATTTCT | 399 |
| Mouse | 286 | GAGAGTAAAGTGACATACCAAGAGCAGATTTTGCGGGCTGAGAACTCTGGGGCTGAGAACTCTGGGGCTGATTTCTAAAGGCCCATPGAGTACTCT | 385 |
| Human | 400 | ACCTTTCTGGAGCTCTTCTGGGAGAGACAAAGATGTTCTCC-TGCCCTAAGACACAGGGACACACACGGGGGTCTGACACGGGCAACACAAATT | 498 |
| Mouse | 386 | ATCTCTTCCTGGGAGGAACTCTGCGGGAGACAAAGATGTTCTCCCTGGGAGAGACACACAGGGACACACACGGGGTTGTAACAGGGSACGGCGAGTA | 485 |
| Human | 499 | GGCGCGGGGAAAGCAAAACACACTGAGCTTGTAGAGCTCCACAAACCTTGTTCTCCAGAGCAGCTCCAGAGCTGGGGGGGGGGGGAGG | 598 |
| Mouse | 486 | GATTCCTCTGGGAGCGAGAGACCCCTCTACCGGAGAGACTGAGCTGAGACGCGGG | 585 |
| Human | 599 | GGCACCCACAGTATGGCTCTCTGTGCGCTTGGAAAAGTTTTTCTACCGGTATGCGCTAAACACGGACACACAGAGAAAGTGACTGTGCACTTAGGGCG | 698 |
| Mouse | 586 | ATTGATGAACTGCTCCCGAAAGTCCCGGAGGTGTGTTCAACTAAATTGGTGTGCTCTGAGTCTC | 653 |
| Human | 699 | CCTGTGTGTACCCGTGTCGTTTGTGGAAACCTCTACAGGCACTCAGGCCGGGGCGCATGGCTACGCCGTGTAATCCAGCACTTGTGGGGCGAGGGGG | 798 |
| Human | 799 | CGATCACCTGAGGTGGGGAGTTCGACACAGGCTGGCAACATGGTGAACCCGGAGGGAGTTGCACTGAGCCGAGATCACACCACTGCACTCCAGCT | 898 |
| Human | 899 | CTGTGATCCCAAGCTACTCGGGAGGTGAGGAGAATCGCTTGAACCCGGAGGGAGTTGCACTGAGCCGAGATCACACCACTGCACTCCAGCT | 998 |
| Human | 999 | GGCGACAAAGAGCGAAATTCCCTCTAAAAAATAAAATGATAATTAAAGCCCATCAACTCACACCTTCAACGGGTTACTGGTGTGTTGAT | 1098 |

FIG. 3C

The Genomic DNA Sequence Homology
Between Human and Mouse *Csx/Nkx2-5*



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FIG. 4A (1)

CTCGAGCCCAGGAGTTCAAGACCAGCCTGGAAACATAGGGAGACCCCTCTCTCTCCACAAAAAAATTAAAAGTAGCCAGGTGGTGGCAAACA
CCTGTAGTCCCAGCTACTCAGAAGGCTGAGGTGGGAGGATCACCTGAG
CCTGGAAAGTAGAGGGCTACAGTGAGCCGTGATCACACCACTGCACCTCC
AGCCTGGGAGACAGAGTGAGACCCCTGTCAAATAAAATAACAAACAAAT
AATGATTAATAACTAAAACATAAATTTATGCTATTTCACCTTGTAT
TTTGTAAAGATTTTAAATGAAAATTCCCAAATTGCTTCCAGAAGG
ATTGTTCAAATTATACCCACATTCACTCATGTTCTCTCCTGAACA
GCAGCAATCAGGAAAAACTCCCTGGAAGAGGGCAGGGCTAGACTGAGA
TTTAAAAGGGGTAGGCCTCAGCTCTCCTCCAGGTTACACTGTGC
ATGTTTCAAACCTCAAAGAATTACACTCTCTGGTTGCATTGCTCTG
TAAAGATCTGACCCACTACTATGTATTAAAAAGGGATGCATGATAATG
AATTCAAGCCCTCTGTAAAATCCAAAGGGCCTATTGAGTTCCCC
CATTAAATGGTCATTAAAATATTCTGGAAAGGACAAAGCTTAGTT
AACTATGAGAAAAACAAGCAGAACAGCCCTGGATTCTGTCTCAAAG
ATTTTACCATGTTGGCAGGCCTGGTAGTCCAGAGCCCAAGAAAATATC
CCAGCCACAGATAACCTAGATGTAGACTAGCAGTGCTACAACCTCAAG
GTCAGAAGTATGTCACTAGACCAGAGCCAAAATAGGTGCTATATCAT
TAAGAGAGTAAAATGCAAACCACAGACAGGGTACATTATTACAAT
AAGCATATAACCCACAGGGACTCCTATCTGAATATGCAAAGAACTCT
CACTAATCAATAAGAAAAGGAAAAGATTAAACAGGCACCTCACAA
AAAAGTATATTCAAATAACATTGAAAAGATCCTCAATT
CACTAGTTATTAGGGAAAGGTGAAATAAAACCACAATGAGACACCCCC
ACGCCACCAGAACGGCTTAAATCTAAAACATGTAATACCGAATG
TTGCAAGGATGCCAGAACACTGCCATTGTACACTGCCAGTATGA
GGGTAAATCTGTACAACCAGGTTGGAAAACGCTGAGTAGAATGTACTC
TAGCTGGATTGTGAATATCATATGATCCAGCAATTCTACTCCTAGAA
ATTTACCCAACAGAAATGTGTAACATGTTACCAAAAGACACACGCA
AGACAATTGATAGAGGCACACTCACTATTCTAACAGTCAAAACTGGAA
ACTACCCAAATGTCATCAGCAGAGAATGGCGATAAACAGTAGCATCT
TCACATAATGAAATGTTGCACAGCAATGAAAAGTAGCTAGCTACAAC
TACAAACAATGTGATTGAAACCTCACAAACATATACTAAGTAAAATTAT
CAGACACAAAGAGTGTATATACTGTATTAGATACTGTGAAGTCTGA
AAACAGGCAAAACTATTCTGTTAGAAGTCAGAATAGTTACTGCC
TGCCGGAAACAGAACTCAAGAGGGCTTAGTAGCTACTGGTAATGTT
TGCTTCTGAACTGCATGCTAGTGAGGCAGCTGTTATTGTGCAGTC
CTGTGTTACACTGGAGTTAAAGTCCCCAAAATCAGAAAGTGTCA
GCAAGTGGAAGCAAGTACACTGCTGGACTTGGCTGGAACTTAGGGGA
TCCCATAATTGTCACAGGCACAAGCAAAGCCAGCTTCTGCCNTAA
GTAGCATCTCCCAGAGTCAGGATCCAGGAATGGTTGGCAGGCAGGAT
GCAAGGCAGGATTGGGAGTGGCTGAGAGTTTCCCAGTGCCACCTGG
TCCCACCTCCCTCTCCCACCTCTAAATGAAACGGGCAGTACAGCTCTG
TTAGGAAAAGAGCCTGGTCCCTAGGCGATGACTGTACATCTAGGGA
GAGGGCGATGCACTGGGTCCTCACCTACACCCCCCTGGCTGTCTCA
CCACTCTGAATTATAATGCCGGACTTCCCTCATCTCCCACCCACACA

FIG. 4A (2)

TCTTGTAGAAGAAAAGAAACGAATCTCCCAGGGCTCCTCTAACAAA
AGTGTTCATTCAAGAGTAGCCCTGCTTGAGGGCCCTGGCCTGGAGGAG
TGGGAGAGGCAGCCCTCCCCCTCCAGGAGAGTCATCTCCAGGGCTACC
CAGGACTGAGTAACTAGGTCAACCAGAGTAACCAAAGAGGCAGGAGACA
AGGGCATTCAAGCATTGGGCCAGGAATGGAGGGTGATGTCCAGTTCAT
GTTCTTCTGGTTCCAGCATAGCACACGGTCAAATGAACCATCATGCA
AGAAAACACAGCTAGTCTCCCTCCACCAGCAACCTTGTTACT
GATAATAATCAAATTCACTATTTTTTTTTAACTAAGGCTGAG
ATAATGTCAAAGGACCACAGGAATAGGAAGGCCTAACCAAGGCCTT
AAAGAATGAGAAGAAGATTCAAAAGCCTCTAAGGGAGGAAG
ATGTTTTCCCTCCTTACTTTCTACAGTAATTTATTTGGATAA
ATAAACCTGATAAATGAGAACCCACGCTTCCCAAGGCCAGGCTGTG
TTTGGTGGGTGGTCCTCCGTCAAGCAGTTGGAGTAATCCAGAGTGATC
CCGGGCAAGTCGAAGGGAGCAAGTCTGTGTGAAGCCAAGAGGTATC
TTCCCTACAGCTCTCAAGAGAGGGATCCCGTGGTAATTGTGAG
GCTGGAAACACCGAGAGGCTGACTCCATGTTATAGAGGTCAATTGAT
GGGTTTGTGCATGGAAGGCAGGAGGAGACTGAGAGTGCTTGTATTG
TTATTTGGTTATTTTATTTAAAAACTGGATCAGCCGACTTGA
ATACAGAAAATGAAAATGAGGAGATTGCATAACAGCGCTGGACGT
CTGAAGGGGCCAGGGCTAGCGGTGGTGGGGCACCTAGAAACACTT
CTGCCTGCAGATCGCGAGGGTTAGCCACAGGAAGGGTGCCTAGGC
TGGCCACAGGGCTTGCTGTGACTGAAGGACCAGCCTGGCGCAC
TTCTTCCCTCTGCCCTGCACTCCGGCCCGCCGGAGTCAGAGCTGA
CTTGCTGCAGGTTGGGGAGAGGACAGAGGCTAGGACGGTGGCGAAACC
TCACCTCGTCGAGTCGGAAAGGTAACCTGGACCCGGCAGGCACCTC
CTAAAGTCCAAGCTGCCCTCTGAAGAATAACCTGATTTCTCCG
GACCGGACAAAGGAGGATCGCTACAACTAGCCTGTAACAAAGATT
CCCTATTTCTGTGGTTAGGAAAAAAAAAAAAAGGAAGCCCTCCGGGA
GAGACATGCGCCCTAATATTTCTCCAGATGGCCGGTTCAAGCGCG
TTTGAGAGTTGCTCTCCTACCAGCCTGGGTCTAGGCCCCCGCAC
CCTCATCTGGCTCCGCCCTCTCCACCCCTCCGGACCCCTAAA
GGGGCGGGGGGCCAAGCCGAGGGCGCTGCGCCTGACCCCGAGCGGA
AGGGCCCCAGTCTAGGTCTTAATGCGGTGGCGTCTCTTGACAGGC
GGCGTTGGGACAACAGCGGGACGAGAGATAAGGTGACATACCAGA
GCAGATTGGTGCAGCGCTGATACTCCTCTCCGACAGGAACCGGG
AGCTATTTAAAAGACCCATCGATTACTTATCTTCTGGAAAGCTT
CTTGCAGAGAGACAAAGATGTTCCCTGCCTAAAGACACAAGGCCACA
CAACGGAGGGTCTGCACAGCGACGCACAATTGGCGGGAAAGCA
AAAACACACTGACGCTTAGAGTCACAAACGTGTGTTCCAGAGCA
GCTCCAGAGTGCAGGGACGCTGGGGCGGGCAGGGCACCCACAG
TATGGTCTCTGTGCCCTGGAAAGTTTTTACCGTATGCGCGTA
AAACACGCACACACAGAGAAAGTGAATGTGCACCTAGGGCGCTGTGT
GTACCCGTGTCGTTAGCGAATTAAAGCACATCAGGCCGGCGCCA
TGGCTCACGCTGTAATCCAGCAGCTTAGGAGGCCAGGCAGGCCGA
TCACCTGAGGTGGGAGTTGACACCAGCCTGGCAACATGGTGAAAC

FIG. 4A (3)

CCTGTCTCTACAAAAAAATACAAAATTAGCCGGCATGGTATGCGTG
CCTGTGATCCCAGCTACTCGGGAGGCTGAGGCAGGAGAACGCTTGAA
CCCAGGGAGGCGGGAGGTTGCAGTGAGCCGAGATCACACCACTGCACTCC
AGCCTGGCGACAAGAGCGAAATTCCGTCTAAAAAAATAAATAAAT
AAAATGATAATTAAGCCCATCAACTCACATTCAAAGCGGTTACTGGTG
GTTGTAATGTATCCATAGACACAGGTCTAAATGTAACGCTCCATTG
TGCTCCTTTAAGGGCTTGAATGTCTGCAACTGTCATGTGTACACTTA
AAGTATGGATGTGCAACACGACCCCTTAGCGCGCTCGTTCGTG
TCTGAATCCCCGCAATTGCCAATTGCTTGGAGCGCAGAACGCCCTC
CGCGAAAGGCGGCTGCTGATCCGACTTTGCTCCGGTATCGCGCAGCT
TGTTGGCCTCCGGTCCCCGTGCCATGCCCGGGAGGCTCCACA
GACACCGCTTGCGCCGAATTATAACGAGACTGAATGGGTTTTGGTG
TGTGTGCAACACAATTGTCAGCTGCTGTTACAATGCGCTCC
GCCGGGCGGTGGAAACTTGGCTGCGGTAACGCACAGCAGGTTGGAGGG
CACGACCCGAAAGGAAGGAAGGAGGGAGGAGGGAAAGGCCGACCCCT
AGGCCCGCTGGCCAGCGTTCCAGCATCAATTGAGCACTGAGCCGGC
CGCAGCAGCACAGGGCTGGGGCTCCCGGAAGTTCGCCAGGCCGGGGT
TTGGGCCAGAGCCCGGGAGGCTGCCGGTGGTAGGTGCGACTCTTCAC
CTCTCCGGGAGCGCGGCCAGCACCCAAACCCACCCGAAGCGCTGC
CGTCGGCCGGCTGGTCCCCCGCGCGGGCACAAAAACAGGCGGCAGTT
CGCCAGCTCTTTCCAAACCTGAACGCCAAGCGGAAGGTTCTTC
CAAAGTCGCGGTTCCCCGGGTTCAACACCCGCCGGCAGGCGCGAAC
AGCCCCAGGACAACCATTTCCTCTCACTGATCTGAGTCGGTGTCC
ATCTGACTCGAATGTCACCTGATTTCAGCTGTGACCTCCAGCGAC
GGGACTCCGAGGAACTGATTCCAGCGTCTCGATTCTCCGCCCTCTCC
GCCCGTTTGGCTGAAGCGGTTGCAGCGTGGGGCAGAAGGGGTGG
GATGTGGCAGCCACCAGCCCCAGCCCAGAGAAGAAAAGAGGACGAAAT
TAACCGCAAAGGACACCGGAAGTCTGAAAGCGACTCCCTCGGATCCTC
GGAATCCGAGGCAAACCTAAACACTAGTTGAAAGCGGATCATATCCA
CTAATCCAGGACAAATTGGGTTGGAAACATACTCCCCAGAGCCTAA
GAAAACGACTTACAACAAAACAAAAGTACAAGGACAAAATGCAAAG
GAGTTTGAAACGTAATTGCTCTCAGAAAATATGTGTATATATATAC
ATCCTATAATATGTTAAATTGAAAAAAAGTCTCTAAGAGGAT
ATATTTTAAACAGTGGCAGCTGGGAGGGAGTGGGATTAGCTGA
GAAGGGGAGAAGGAAGCATTGAGGTGACGTAAATGTTTGAT
TTGATTATGGTGGCTGTTATGGGGGTGCACATCCAAGTGTCAAGACTC
ATCGAAGTACACTTTGTTCTAGGTACATTAGACCTCAATAAGTG
GATTTAAACCTAAATAAGCCAGGTAAACAGCTTGCTGGTGGCTGG
GGGAGAGGCTGGGACACTTACATTGATCTCCCTCTAGGCATGTT
GTTTGGTTGGTTGTTATGATGTATTATTATTCAAAAATAT
ATCATTAGCAGAGTGACTGATGTAATGTAACACCATTGTTAAGGAAA
CCAACAAAAGCGGGAAACAAGAGACACTGGTGCATCCTGTTAGAGGGAT
AAGAATAAGCACTCGCTGTCAGCTCATAAAATTTGGGAATGAA
TGTGTTCCGCTTGTGTTGGTTGGCTCATGTGTTAACAT
CAACGAGAAATGAGGACCCAAAATTATCCAGTGGTACGTGTTG

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FIG. 4A (4)

GTGTGGCTGTCATCTCCTGGACTGGCTACTGAAGGCCACAGGCCTG
GGAGGACCAAATGCTCCCTGGATGTTGAGTCCCAGCCGGTAAGCAGCA
CACAGTCCCCTGAGCAAAGATGTGGTGGCCGGCTCGCGCTGTGGGG
GAAGGCCAGGCCGGACAGGAACCTCAGATCTCACCGGCGGATGAGAG
TGGTGCCCCCTGAGCTGGAGTCCCTGCTGGCCTGAGAGCTCAGCTG
TGCCACCGTTGGCAGACCCCCACACTTCAGGGAGCTGCCAGGATCAGT
GGCTACAAGAGTCCCCACCGTGGAGAAACTAGGTATGAAATATT
TCCATTTACACCCCTACCCCGCCCCAGACAGGAAAGTCACTTCAACC
TTGTTAGGTAGATTCCAGATCTGGTCAGATGCAGGGCTATTCAGA
GAGATTTTAGAGGCTGACTCTCAGGAGAGGAAAGGACAGTGGCTGA
AGGCCAGGGTCAGGAAATCTAGGAAC TGCTAAACTCCTCTGCTGGCC
TGCAGGGAGGCCCGGGTGGGCTACCAAGGCCAACAGGCAAGTCCAT
CTTCCCACCTTGCCACCTCTCACAGGGACCAGGCTCTGCATCCTCAG
TGACCAACAAGACTGGGCCTGCCCTAGTTGTCTATACTGCC
TCCCTTGACTCATACTGTCCAAGACCCAAGACCAAACCAAGTCAG
GAGAGATTTGAGGGCAGCCAGTGCACCCAGGGCTGTTCCAGGTA
CTACTAGACAAAGGCCACCCCTCCCTCTCTAGGGCTCCGCTG
ACCACCTGCACAGTCTCCTACACCAAGGGCTCCGGTGCACCCCTT
CACAGAGAGTTCACTGCACCGCTGCTCGGCTGCCTGTCTCAAACCAT
ACACACACCTTGATTCTAAACTCCAAGATTAGGATGGCCCCAGAA
ATCTGCATTTTAATATGTACCTCAGAGGATTCTGGCCTAGATATTTC
TACAGCCCCAAAAGTAACAAGGAACCTGTTCCAAAAAGTGTATTACGG
AAACTGTCATGTTATTCTGACTTGCCCCCAATTATTCTCCCTG
AAGTTTCATCACCAAAAAACCCACATGTGAACCATATGTGTACATA
TGCCCCATATTAAAATACAAATTCTGCACCTGGTTGCTATTAAAGT
ATCTCAAAACATATCCATAAGAATACATATGAATGGAACTAATTCTT
CTCATGGGATATGGGATCTGTTCTATGGACAAACATAATTAAACCAG
TCCTAGTATATACACTGGTTTACATGTTGATCTTAAAAATAA
AACGGNTGAAA (SEQ ID NO.: 4)

FIG. 4B (1)

CAATTTCTATTNAGTCTATTAAAAGGGATTTTTTNAACTCACTGGNAACCAGGAGGA
CTGNAAAGAAAAGTGAATGGCTCTGGACTTCCTCTAAGGAGACCAGCATGGTCGCC
CCAATTTATTTGCACGTATTGTCCTGGCCCATCTCCTCTCCTGAAACAC
CAAGACCTTTGGAAGCCAAGAGAAATCATTACCCGATTCAAAAGAGCATAGAGAGTG
TAACAGTCACTGATCTTGTCAAATAGGGAGAGTTTTCTCCCTTTGTAACAC
CTGACCCACAGGACTGACAGTTCTAGGAAGCCCCCTACCGAAAATAGGAAATAATCC
TTGCCACCTTGATTGCAAGGGCAATGCTAATTTCTTCTCCAGAGCTCTCAAAAA
AAAAAAAAAAACCTTACTAAAAACAGGGATCCGGATGTAGCCTCGATGTCCCCAT
TAAACGGTAATATTCAGGCGTCCGCTCACACTAATCTTCAAACGTCTCGAGGCCG
CCTGGCCAGCAGATTCACTAACAGCGCTCCAGGACCCTCGTCCGAGCTTTTCAGC
GAGACATTAAATTGAATCGGATGTGGCTCGTTGCCAGACGTACCGCCTCGCGATAGG
CATCCTCTCCAACGACACCCCCCCCAGCGCTCGAAAACAATCTTAAAAGGCAAGG
GGGCCCCCAAGTAGGTTAATTACAAACATAACGGTAACGTGGCCAAAGNCAGGCGAG
GAAGGGCCGCAAGGCCGCTGACATGCAAGCTCCGTCCAAGAAGAATTGGGTTGGAGGTG
AAGAGGTGGGGGAGCAGGTTCTGGGCTTGAACGCCCCACATTAAAAAGGCATCC
TCCACAGACTAGACTAACAAATTCCAGACCCCCAGTAGTCCCTGGCTCAGAAACTCGAGGC
GTGATTCGGCGTGGCAGCCCAGGCCTGTTACTGACGGCTGGCGCTAGAAGCCGGGTC
AGGGCGTTGCGCGCCTCTGGCTGCCCTGCGGGCTCACCTCTCCCCAGCATGGAGG
CCCCAGGTCTGGAGTGTGGCTTGTAGGAGACAGGAAAGTCCAACATCAGGCCAA
TGCTTGACTTCACTTGCCTGGCGTCTCAGACGGCACACTGTGGGTTGAGCACCCAAAG
ATGTACGTTCTGGACAGACACTATTGTCCCCATACATGGAGCGTTCTCCGCACCTT
GGCGCGCCTGCGGGAGCTGTCTTAGGTAGTTTGCCCTGCGCCGCTTATTCT
ACTCCAAGCGCTTTGCCAACCCGCACTCCGAAAGAGCCAAGCCCTCCACATCCCCA
TTCTCAGCAAGTCCACGCGTCCGCCAGCTCCGCCCGGGTCCCTGTACCAAGCTAG
GGCGTGAGAAGCCAACGCTTCACTGACAATCCTGTATCCCCAGCTAGAAGGC
GTCCTTAACCTGGGCCGCTCTGCCCTGCCGGACTCCTGAATTGTAAGAAAATAAAACT
CCTCTCTGCACTGGTCTGGGAATGGAGAAGACCCCAAGCTTCATCAGACCCCTCCAAAG
GAGTGCAGGGGACCCAGAGAAATGAGGCCACCCGGGAGGATCTGGCAATGTAGCTGGCGC
TCCTGAAACTCTGGCAGATTGTCTGACTTCTGTGCCCTACTCTACTGACCCCTGGGCTAA
AAATGATCATGATCACCCACTTGCCTGCCCTCCCCACGCGCCTGACCGAGCCGCAG
GGGTGCCCACTGGAAGTCCGCCAGAGGCCTCAGAGAAATCCTGGCTAGCTGGGCTC
AGAGGAGCCCCGCCCTCGAGAGCTAAACCTGGCTAGGACCCCTGAAACCTCGAGGTTG
GCAGAAGCCTGAGGGCTTGTGCCAGGCAGGGAGGGCACGGGAAGGAGGGAGGTGGGAT
CGATGGCCTCCAAACAGGGAAACAAGGTGGCTGGTAGCTGGGCACTCCACAAGACAGG
TGTNTCCTGGGAAGCTGAGTTACCAAGCTGGATTCTGATTTCATTATTAAGGGG
AGAGGCATTCCCTGGGAGGGTACTGGCAGTGACTGATGCCCTGGAGTTGTGCTGTG
CATAACACTACTGTAGGAGGCAGCAACTCCTACCCACCTGGCATCACTCACCTGCC
TTACTTCTGTTGATTGCCAGAACGACCCAGAGCCTGCCATGATTGACCCCTGTAGGC
CAAGCCAAACCAACCCGAATTGTCAGAATTTCGCCCTGGTAGTCCCCAAAGCCC
AGCCCTGTCTTNAGGTTTTCTATTGAGATTCCCTCATCCACCCACCTTAGT
AATAAAGCCTTCCTCAAACTAATTCTCCCCACCGCTCCACCCATCCTTTTTTT
CCCATGCTGGTTGGGTGCTGAGGAATTCTCAAACCCACCCATCCAGCCCTGCC
CAGAGGCCCTGACTTGCATGCCCTGGTAGGNTTCAAGGGTTACATTAGGGAGCAAAAG
CAGGGTGCAGGGCAAAAGGGACCCCTCCAAATGGTCGTGCCCTTAAAAAAGCTG
GGCAGGGNTTTTTTTTTTTTTTTTTTTGCGTATGACTATA

FIG. 4B (2)

TTAGGTGACACGAAACTGCTCATCGCTCCTGTATCGAGGCCCTGGCCAATGGCAGGC
TGAGTCCCCCTCCCTGGCCTGGTCCCGCCTCTGCCCTGTGCTCAGCGCTACCTG
CTGCCCAGACACATCCAGAGCTGGCGACGGGTGCGCGGGCGGGCAGGACCATGCAG
GGAAGCTGCCAGGGCCGTGGCAGCGCCGCTTCTGCCGCCACCTGGCGCTGTGAGAC
TGGCGCTGCCACCATGTTCCCAGGCCCTGCTCTCACGCCACGCCCTTCAGTCAAAGA
CATCCTAAACCTGGAACACAGCAGCAGCGCAGCCTGGCTGCCGCCGGAGAGCTCTGCCCG
CCTGGAGGCAGCCCTGGCGCCCTCTGCATGCTGGCGCCTCAAGCCAGAGGCCA
CGCTGGGCCGAGGCCGTGCGCCGGCCTCCAGAGCTGCGCGCAGAGCTGGGCCGCG
GCCCTCACCGCCAAGTGTGCGTCTGCCCTTCCGCCGCCCTATCCACGTGC
CTACAGCGACCCCCGACCCAGCCAAGGACCCTAGAGCCGAAAAGAAAGGTGAGGAGGAAAC
ACAGGCCCTCTCCCTGGTCGCTTCGTCCCCAAGAAACTCAGGCCAGGAGG
AGGACACACGCCCTGGCCGAGGGCTGGCTGCCGGGGGTCAGAATGTAAGAT
GCCTGGTGTGCGCCAGGCTCCCGGCCGCTCCAATCGAGGTTCAAGAGGAAATGC
CGGATTGAAAGGATCCGAAAGCAAGAGACCAAAAAACTTTCCCCCGCCCTAACAAACC
CCCGCGGTTCCGCTCTGCTCTGGTTCTGGTAGAATTAAAATGGTTATGGTTA
AACAAAACAAAAAACAGCCAACCCCCCTGGATTTACCCCCCTTGGATTTCAAACC
CTTTTAAAATTTGAAAAAAACCCCCAACAAAATTAAATTTTCCCCCAAAAT
TTTTTTTTAACAAAAGGGGGGTTGGAAAATTTTTCCCCCCCCAAAGGGGTT
TTTGTTTTTTTT-----TTTNTTGGCAAAATGAATTNTGGANCAGGCCTTAT
TTNAAATGGATATTGGNCCNCAGGATTTGATTCATTATTTTAAGCAAACCTTNC
CGGGCCGGCAAGGGAAAGGTTCCCTCGTGGAAAAGTAGGAAATGCTGCGCTACCGCGGG
ACAAGGNAGTGGACGAGATGAGTGCAGGATCATCCCGCAGGCCATCCAGGATGGGGA
GGGAGGCCGGCCCCGCTGCAGAAAGGGCTCTGGAGACCCCCCAGCCAAGGCAGGAG
CCCGGGCGATTCCGGAGGCCGAGCGCTGGCGAAGCGCTGGCGAAGGGCCGCTGC
CAGCCGGAGAGAATTCAAGGTTGTTGAGGAGCAGAGGCTGGAAACAAATTGGGCG
GGCACGGCGCTAGAACTGATCGCTACCAATTGAGGAAGCCAGCAAGGCAGGTTCCGAG
GCCGCTGCCACCCGAGCTTGGACACTGCGAAACCTGCTGCCGCCAGGCTGGA
GCCTCCGATCACCAAACACACTCCCTGGCTTCTGTTCTGATTCTTAATTGAG
ATAAGACCGTCCCTAGCAGTGAGGCTCGGCCTCTGTCATTAACTCTCAAACCAAAC
TAGCCCTAATTCAAGTTCACCCAGAGCATCACCTGGTTTATTAAATTTTTATT
TTATTATTTTTTTTGTGAGGCTGAAATTAAAGTCACCGTTGTCTCCCTCACC
AGGGTGTGAACGTCCCCGAGGGCAGAGACCTCCGTTGTTCCAGCGCTTGAGCCA
GCTTGACTTTTACAATGCTGAGTGAGACGTGCGTGGCTCCAGTGCACCTGGCAGA
GTGAGGCCAGCCAGCTGGCGCTCCAGGACAGTCAGTGGCTCCACGAGGATCCCTT
ACCATTACTGTGCGGCCGCGCTCCGTAGGTCAAGCCCTTACCAAGCGTCTTCTGCC
TTTCTGTTCCCCCTCAGAGCTGTGCGCGCTGAGAAGGCCGGTGGAGCTGGAGAAAGACAGA
GGCGGACAACGCCGAGCGGCCCCGGCGCAGGGCGAGGAAGGCCGCGTGCCTTCTC
GCAGGCCAGGTCTATGAGCTGGAGCGCGCTCAAGCAGCAGCGTACCTGTCGGCCCG
CGAACGCCGACCAGCTGGCCAGCGTGAACACTCACGTCCACCGCAGGTCAAGATCTGGTT
CCAGAACGGCGCTACAAGTCAAGCGCAGCGCAGGACCAAGACTCTGGAGCTGGTGG
GCTGCCCGCCGCCGCCGCTGCCAGGATCGGGTGCAGTGCCTGGTGCAG
TGGCAAGCCATGCCCTAGGGACTCGGCCCTACGCCCTGCCACGGCGTGGCCTCAA
TCCCTACGGTTATAACGCCCTACCCGCCATCCGGTTACGGCGCGGGCTGCAGGCC
TGGCTACAGCTGCACTGCCGCTACCCGCCGCTTCCCAAGCGCAGCCGGCACTGC
CGCCGCCAACAAACACTCGTGAACCTCGGCCGTCGGGACTTGAATGCGGTTAGAGCCC

FIG. 4B (3)

CGGGATTCCGCAGAGCAACTCGGGAGTGTCCACGCTGCATGGTATCCGAGCCTGGTAGGG
AAGGGACCCCGCGTGGCGCACCTGACCGATCCCACCTCAACAGCTCCCTGACTCTCGTG
GGGAGAAGGGGCTCCAACATGACCCTGAGTCCCCTGGATTTGCATTCACTCCTGCGGA
GACCTAGGAACCTTTCTGCCCCACCGCGTTGTTCTGCGCACGGGAGAGTTGTGGC
GGCGATTATGCAGCGTGCAATGAGTGATCCTGCAGCCTGGTGTCTAGCTGTCCCCCAG
GAGTGCCCTCCGAGAGTCCATGGCACCCCCGGTTGGAACGGAACTGAGCTCGGGCACG
CAGGGCCTGAGATCTGGCCGCCATTCCGCAGCCAGGGCCGGCGCCGGCCTTGCT
ATCTCGCCGTCGCCCCACGCACCCACCGTATTTATGTTTACCTATTGCTGTAAG
AAATGACGATCCCTTCCATTAAAGAGAGTGCCTGACCCGCACGTGTGCTTCTTCA
GCTTGCAGCGCTTCAGAACAGCAGGAGAGAGGTGGCCGCCGGACTGGTCTCAGATCTCAG
GCACAGGATTCCCTGAGCAAATTGATAACATTGATAACTAATAAAACCTAACCTTGCTG
GAACCATACTGGTCCGTGCGGCACTTCTGAGATTGTCTCATATAATCCTCAATAAT
CCAACAAAAAAATCTAAAGTTAGAACAGCTGAGGCCGGAGAGGTTAATGACTTAC
CTGCGAGCAAATAGCCAGTACTAGTCGAACTCTGGTTAAATTAGGATGCCTCACTTCAG
AGACCGCCTCCCTGTGCTCCCAAGCTCCCCCTTGAATCCTAATGTGTGCCAGGCACG
GTTCCAGGCACTGGCATTAAATGGACAAGCAAAAGAACCTGGCCCTGTAGCTGGAG
AGCACCGTGATCATCCCACTAAAGAACCTCTAACCTGTTCCAAGATGGNAAAAGCC
AAGAANCCAAGCCCTGGNAAGCGTTCTCAAGGGCCTCANATGCCCAAATGCCACG
TCGGGGGCTCAACANCTNGCCGTTGAACTGAATGCCNANGGTGGCCCCAAANAAGGN
TCCTGCGGGATGGNGCTCAACTCCAAGCTGTGGTGAAGGCCATAAAATTCAAATGGGCC
AAGGGGAGCCCCCTAAAGCCTAAACCTCNGGGGTCCNTCCCTAAGGGATTTAANT
TTACCAAAAGTTGGNCAAANAATGTTCCAATGGNCNGATTATNGANGGGNAAAAC
TGGNGGGCAACCGAAATCCAGTTAAACCCGGTTGTT (SEQ ID NO.: 5)

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FIG. 5A

AGGCCCGCCG CACCCCTCATC CTGGCTCCCG CCCCTTCTCT CCACCCCTCCC
GGACCCCTAA AGGGCGGCG GGGCCAAGC CGAGGGCGCT GCGCCTGACC
CCGAGCGGAA GGGCCCCAGT CTAGGTCTTA ATGCGGGTGG CGTCTCCTTT
GACAGGCGGC GTTTGGGGAC AACAGCGGGG ACGAGAGATA AGGTGACATA
CCAGAGCAGA TTTGGTGCAGC GCGCTGATAC TCCTCTCCCG ACAGGAAACG
CGGAGCTATT TAAAAGACCC TATCGATTAC TTTATCTTTC CTGGAAAGCT
TCTTGCGGAG AGACAAAAGA TGTTCCCTGC CTAAAGACAC AAGGCCACAC
AACGGAGGGT CTGCACAGGC GACGC (SEQ ID NO.: 1)

TGCTCCTTT TAAGGGCTTG AATGTCTGCA ACTGTATGT GTACACTTAA
AG (SEQ ID NO.: 2)

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FIG. 5B

AGGCCCGCCG CACCCCTCATC CTGGCTCCCG CCCCTTCTCT CCACCCCTCCC
GGACCCCTAA AGGGGCGGCG GGGCCCAAGC CGAGGGCGCT GCGCCTGACC
CCGAGCGGAA GGGCCCCAGT CTAGGTCTA ATGCGGGTGG CGTCTCCITT
GACAGGCGGC GTTTGGGGAC AACAGCGGGG ACGAGAGATA AGGTGACATA
CCAGAGCAGA TTTGGTGCAC GCGCTGATAC TCCTCTCCCG ACAGGAAACG
CGGAGCTATT TAAAAGACCC TATCGATTAC TTTATCTTC CTGGAAAGCT
TCTTGCAGGAG AGACAAAAGA TGTTCCCTGC CTAAAGACAC AAGGCCACAC
AACGGAGGGT CTGCACAGGC GACGCACAAT TCGGCGCGGG GAAAGCAAAA
ACACACTGAC GCTTAGAGTG CACAAACGTG TGTGTTCCA GAGCAGCTCC
AGAGTGCAGGC AGGGACGCTG GGGGCGGCGA GGGGCACCCA CAGTATGGTC
TTCTGTGCCCT TTGGAAAGTT TTTTTCAACC GTATGCGCGT AAAACACGCA
CACACAGAGA AAGTGAATGTG GCACTTAGGG CGCCTGTGTG TACCCGTGTC
GTTTAGCGA ATTTAAAGCA CATCAGGCCG GGCGCCATGG CTCACGCCTG
TAATCCCAGC ACTTTAGGAG GCCGAGGCGG GCCGATCACC TGAGGTCGGG
AGTTGACAC CAGCCTGGCC AACATGGTGA AACCTGTCT CTACAAAAAA
TACAAAAATT AGCCGGGCAT GGTGATGCGT GCCTGTGATC CCAGCTACTC
GGGAGGCTGA GGCAGGAGAA TCGCTTGAAC CCGGGAGGCG GAGGTTGCAG
TGAGCCGAGA TCACACCACT GCACTCCAGC CTGGGCGACA AGAGCGAAAT
TCCGTCTAAA AAAATAAAAT AAAATAAAAT GATAATTAAG CCCATCAACT
CACATTCAA GCGGTTACTG GTGGTTGTAA TGTATCCATA GACACAGGTC
TAAAATGTAA ACGCTCCATT GTGCTCCTT TAAGGGCTTG AATGTCTGCA
ACTGTCATGT GTACACTTAA AG (SEQ ID NO.: 3)

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FIG. 5C

AGAGAAATCA TTACCCGATT CACAAAGAGC ATAGAGAGTG TAACAGTCAC
TGATCTTGT TCAAATAGGGA GAGTTTTTT TCCTTCCTT TTTGTAACAC
CTGACCCACA GGACTGACAG TTCTAGGAAG CCCCCTTACC CGAAAATAGG
AAATAAAATCC TTGCCACCTT GATTTGCAAG GGCAATGCTA ATTTTTTCT
TTCTCCAGAG CTCTCAAAAA AAAAAAAA AAAACCTTAC TAAAAACAGG
GATCCCAGAT GTAGCCTCGA TGTCCCCAT TAAACGGTAA TATTCAGGC
GTCCGCTCAC ACTAATCTT CAAACTGTCA TCGCGAGCCG CCTGGCCAGC
AGATTCACTT AACAGCGCTC CCAGGACCT CGTTCCGAGC TCTTTCAAGC
GAGACATTAA ATTGAATCGG ATGTGGCTCG TTTGCCAGAC GTCACCGCCT
CGGCGATAGG CATCCTCTCC AACGACAC (SEQ ID NO.: 6)

FIG. 6 Transgenic Constructs of the Human *Csx/Nkx2-5* Enhancer

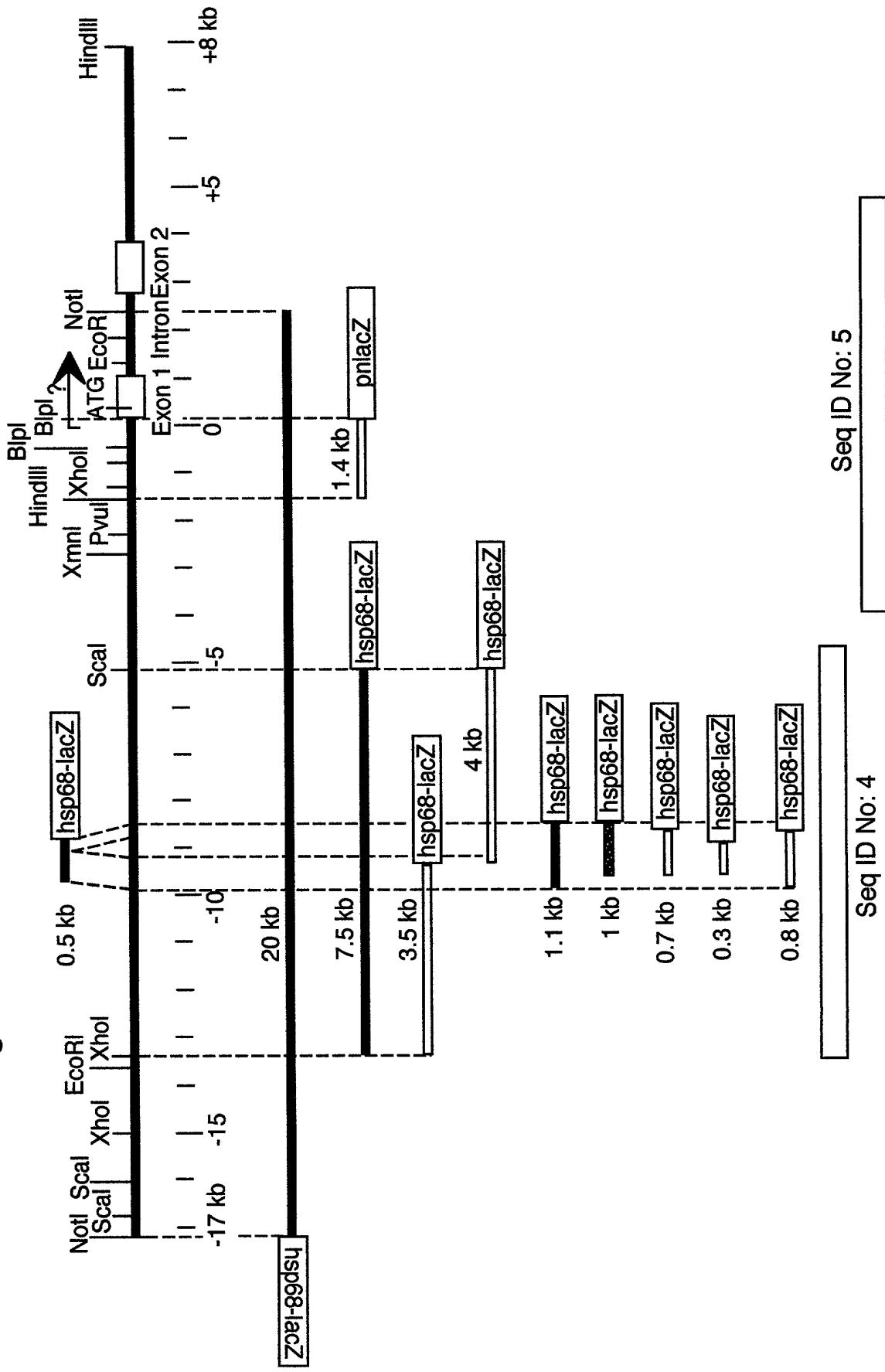


FIG. 7 Transgenic Analysis of the Human Csx Enhancer Sequence

| Constructs | # of Transgenes | Enhancer positives (Cardiac : Ectopic) ¹ |
|------------------------|-----------------|--|
| 20 kb | 8 | 4 : 0 |
| 7.5 kb | 8 | 6 : 1 |
| promoter-proximal 4 kb | 7 | 0 : 1 |
| promoter-distal 3.5 kb | 6 | 0 : 0 |
| 1.1 kb | 8 | 3 : 1 |
| 1.0 kb | 10 | 1 : 2 |
| 0.7 kb | 8 | 0 : 3 |
| 0.3 kb | 11 | 0 : 6 |
| 0.8 kb | 6 | 0 : 1 |
| 0.5 kb | 2 | 2 : 0 |

1. Each embryo was classified into either 'cardiac' or 'ectopic', judged upon the extent of similar to the endogenous Csx expression pattern.

FIG. 8

Cardiac Expression of the hCsx Enhancer-hsp68-lacZ Constructs

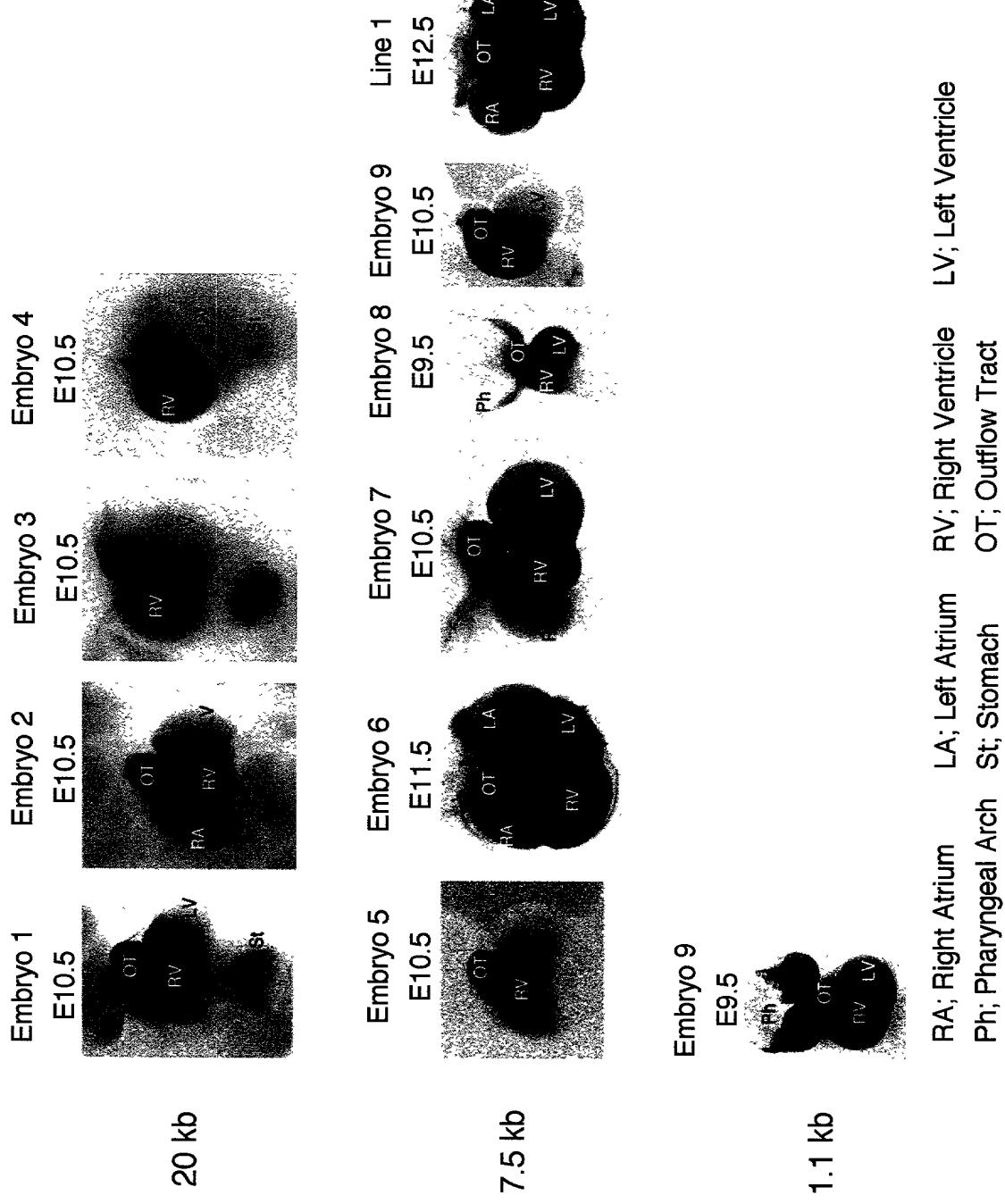


FIG. 9

Cardiac Expression of the 7.5 kb hCsx Enhancer-hsp68-lacZ Construct

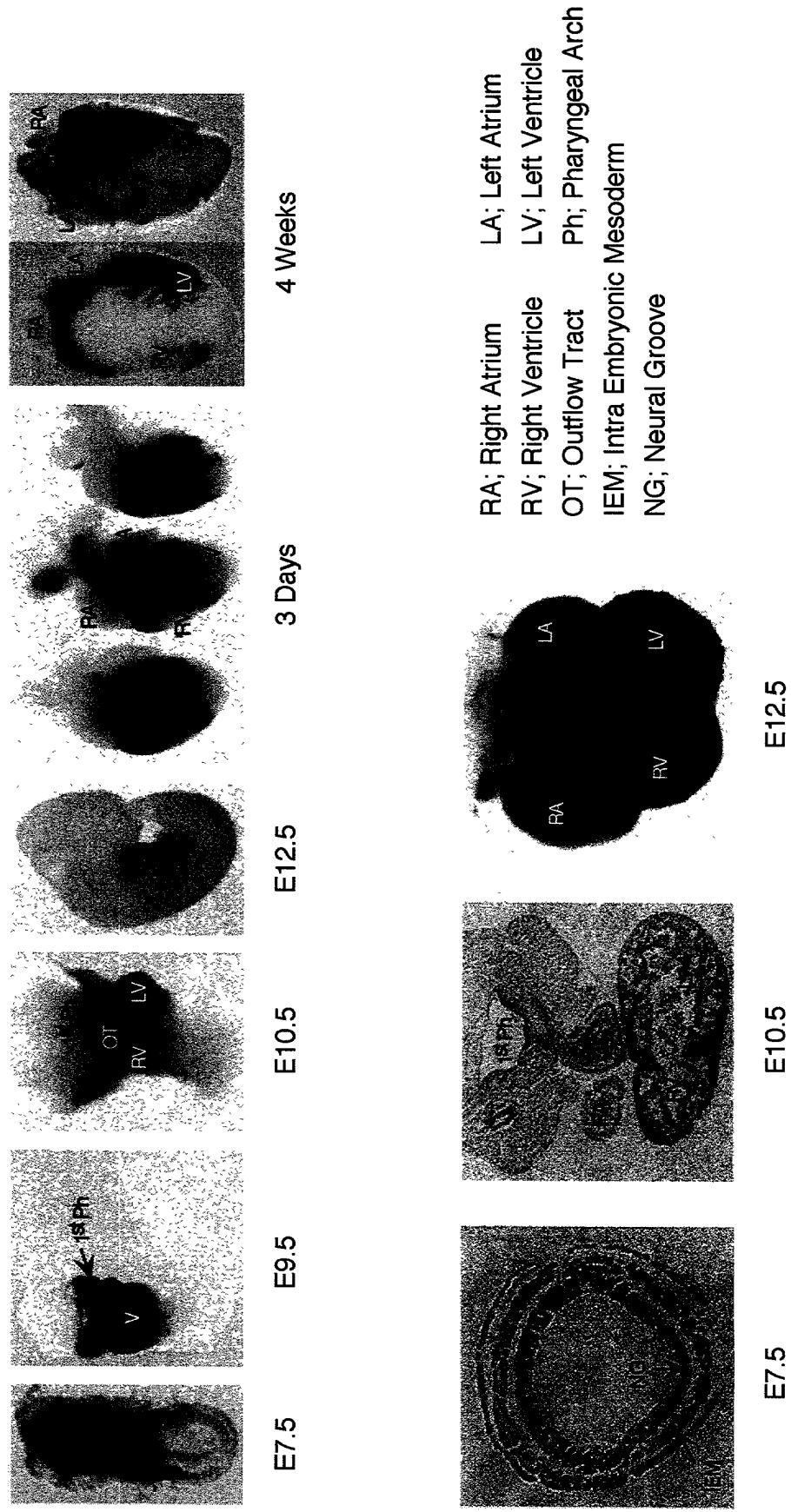


FIG. 10

Facilitated isolation of cardiac myocytes

